

SEQUENCE LISTING

<110> ARES TRADING S.A.

<120> IL-8-LIKE PROTEINS

<130> P032741WO

<140> PCT/GB03/05621

<141> 2003-12-19

<150> GB 0229854.5

<151> 2002-12-20

<160> 24

<170> SeqWin99, version 1.02

<210> 1

<211> 126

<212> DNA

<213> Homo sapiens

<400> 1

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aaatactttc aaagacatga acaatcaaaa cctcatcaag aagaaataga caacctgcat 120
agccct 126

<210> 2

<211> 42

<212> PRT

<213> Homo sapiens

<400> 2

Met Ser Ala Gln His Gly Leu Val Ser Lys Phe Gly Leu Gly Leu Leu
1 5 10 15

Leu Leu Gly Asp Lys Tyr Phe Gln Arg His Glu Gln Ser Lys Pro His
20 25 30

Gln Glu Glu Ile Asp Asn Leu His Ser Pro
35 40

<210> 3

<211> 330

<212> DNA

<213> Homo sapiens

<400> 3

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aggcctcaaa ctgcccccaa agcaggcaaa ggtcagatgt gtggagagag gatggcgagg 180
atggcaagga cggccaagga gggtcgcccc aggtgcctgg acccaggtt gtcccgacc 240
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tgggctcctg gtggcactgg ctggatgctg 330

<210> 4
 <211> 110
 <212> PRT
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<400> 4
 Asp Leu Pro Thr Pro Gly His Pro Val Thr Leu His Ser Leu Cys Phe
 1 5 10 15

Cys Ser Pro Arg Gly Thr Leu Leu Glu Gly Pro Met Ser Ser Gly Phe
 20 25 30

His Arg Phe Glu Val Glu Asn Leu Arg Pro Gln Thr Ala Pro Lys Ala
 35 40 45

Gly Lys Gly Gln Met Cys Gly Glu Arg Met Ala Arg Met Ala Arg Thr
 50 55 60

Ala Lys Glu Gly Arg Pro Arg Cys Leu Asp Pro Gly Leu Ser Arg Thr
 65 70 75 80

Pro His Pro Gly Pro His Val Phe Leu Pro His Ser Pro Thr Pro Ala
 85 90 95

Ser Trp His Gln Trp Ala Pro Gly Gly Thr Gly Trp Met Leu
 100 105 110

<210> 5
 <211> 459
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 <213> Homo sapiens

<400> 5
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 agccctgacc tgccccacgccc gggacaccccg gtgacactcc actccctctg cttttgcaggc 180
 cccccggggga cccttcctcgaa gggccccatg tcttctgggt tccatcgctt tgaggttagaa 240
 aatctgaggc ctcaaactgc ccccaaagca ggcaaaaggc agatgtgtgg agagaggatg 300
 gcgaggatgg caaggacggc caaggagggt cgccccagggt gcctggaccc aggttgtcc 360
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 caccagtggg ctcctggtgg cactggctgg atgctgttag 459

<210> 6
 <211> 152
 <212> PRT
 <213> Homo sapiens

<400> 6
 Met Ser Ala Gln His Gly Leu Val Ser Lys Phe Gly Leu Gly Leu Leu
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Leu Leu Gly Asp Lys Tyr Phe Gln Arg His Glu Gln Ser Lys Pro His
 20 25 30

Gln Glu Glu Ile Asp Asn Leu His Ser Pro Asp Leu Pro Thr Pro Gly
 35 40 45

His Pro Val Thr Leu His Ser Leu Cys Phe Cys Ser Pro Arg Gly Thr
 50 55 60

Leu Leu Glu Gly Pro Met Ser Ser Gly Phe His Arg Phe Glu Val Glu
 65 70 75 80

Asn Leu Arg Pro Gln Thr Ala Pro Lys Ala Gly Lys Gly Gln Met Cys
 85 90 95

Gly Glu Arg Met Ala Arg Met Ala Arg Thr Ala Lys Glu Gly Arg Pro
 100 105 110

Arg Cys Leu Asp Pro Gly Leu Ser Arg Thr Pro His Pro Gly Pro His
 115 120 125

Val Phe Leu Pro His Ser Pro Thr Pro Ala Ser Trp His Gln Trp Ala
 130 135 140

Pro Gly Gly Thr Gly Trp Met Leu
 145 150

<210> 7

<211> 360

<212> DNA

<213> Homo sapiens

<400> 7

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agaataaaaga agtttagtaga tggccttgag tttcccaa caatggcatt ttctgctacc	120
aaaataaaata tggatttcag tcagaaccac tggactataa gaagtatatt ccattctgg	180
tttactggg ggaaaggatg ttgccacaag atgtcagtcc atttattcat tcatatatcc	240
aatagatatt ttatgaccac ttccatgtgc caggagatgg ctaagatct tggaaagacag	300
ataaaatgct acctaccaac tcaaagtcca gtttagggagt caggggtaa aacaatattc	360

<210> 8

<211> 120

<212> PRT

<213> Homo sapiens

<400> 8

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1 5 10 15	

His Asp Asn Ala Arg Ile Lys Lys Leu Val Asp Gly Leu Glu Phe Ser	
20 25 30	

Gln Thr Met Ala Phe Ser Ala Thr Lys Ile Asn Met Leu Phe Ser Gln	
35 40 45	

Asn His Trp Thr Ile Arg Ser Ile Phe His Ser Gly Phe Tyr Trp Gly	
50 55 60	

Lys Gly Cys Cys His Lys Met Ser Val His Leu Phe Ile His Ile Ser
 65 70 75 80

Asn Arg Tyr Phe Met Thr Thr Ser Met Cys Gln Glu Met Ala Lys Ile
 85 90 95

Leu Gly Arg Gln Ile Lys Cys Tyr Leu Pro Thr Gln Ser Pro Val Arg
 100 105 110

Glu Ser Gly Gly Lys Thr Ile Phe
 115 120

<210> 9
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 aaaataaaata tgttatttcag tcagaaccac tggactataa gaagtatatt ccattctgg 180
 ttttactggg ggaaaggatg ttgccacaag atgtcagtcc atttattcat tcatastatcc 240
 aatagatatt ttatgaccac ttccatgtgc caggagatgg ctaagatcct tggaaagacag 300
 ataaaatgct acctaccaac tcaaagtcca gtttagggagt caggggtaa aacaatattc 360

<210> 10
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<400> 10
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 1 5 10 15

His Asp Asn Ala Arg Ile Lys Lys Leu Val Asp Gly Leu Glu Phe Ser
 20 25 30

Gln Thr Met Ala Phe Ser Ala Thr Lys Ile Asn Met Leu Phe Ser Gln
 35 40 45

Asn His Trp Thr Ile Arg Ser Ile Phe His Ser Gly Phe Tyr Trp Gly
 50 55 60

Lys Gly Cys Cys His Lys Met Ser Val His Leu Phe Ile His Ile Ser
 65 70 75 80

Asn Arg Tyr Phe Met Thr Thr Ser Met Cys Gln Glu Met Ala Lys Ile
 85 90 95

Leu Gly Arg Gln Ile Lys Cys Tyr Leu Pro Thr Gln Ser Pro Val Arg
 100 105 110

Glu Ser Gly Gly Lys Thr Ile Phe
 115 120

<210> 11
 <211> 265
 <212> DNA
 <213> Homo sapiens

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 ttattcattc atatatccaa tagatatttt atgaccactt ccatgtgcca ggagatggct 180
 aagatccttg gaagacagat aaaatgctac ctaccaactc aaagtccagt tagggagtca 240
 ggggtaaaa caatattcta gcaca 265

<210> 12
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 12
 Met Ala Phe Ser Ala Thr Lys Ile Asn Met Leu Phe Ser Gln Asn His
 1 5 10 15

Trp Thr Ile Arg Ser Ile Phe His Ser Gly Phe Tyr Trp Gly Lys Gly
 20 25 30

Cys Cys His Lys Met Ser Val His Leu Phe Ile His Ile Ser Asn Arg
 35 40 45

Tyr Phe Met Thr Thr Ser Met Cys Gln Glu Met Ala Lys Ile Leu Gly
 50 55 60

Arg Gln Ile Lys Cys Tyr Leu Pro Thr Gln Ser Pro Val Arg Glu Ser
 65 70 75 80

Gly Gly Lys Thr Ile Phe
 85

<210> 13
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer INSP094-CP1

<400> 13
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<210> 14
 <211> 22
 <212> DNA
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<220>
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<400> 14	
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<211> 37	
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<223> Primer INSP094-EX1	
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<210> 16	
<211> 35	
<212> DNA	
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<220>	
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<400> 16	
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<210> 17	
<211> 37	
<212> DNA	
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ggggacaagt ttgtacaaaa aaggcaggctt cgccacc	37
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<211> 51	
<212> DNA	
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<223> Primer GCP Reverse	
<400> 18	
ggggaccact ttgtacaaga aagctgggtt tcaatggtaa tggtgatgg	51
<210> 19	
<211> 20	
<212> DNA	
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<400> 19	
gccagcttgg cacttgatgt	20
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<211> 20	
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<220>	
<223> Primer pEAK12R	
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gatggaggtg gacgtgtcag	20
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<211> 18	
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<223> Primer 21M13	
<400> 21	
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<210> 22	
<211> 18	
<212> DNA	
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<223> Primer M13REV	
<400> 22	
caggaaacag ctatgacc	18
<210> 23	
<211> 19	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer T7	
<400> 23	
taatacgact cactatagg	19
<210> 24	
<211> 18	
<212> DNA	
<213> Artificial Sequence	
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<223> Primer T3	

<400> 24
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18